Hai Yan Fan

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7480246/publications.pdf

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		1163117	1125743	
15	181	8	13	
papers	citations	h-index	g-index	
15	15	15	144	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	Citations
1	Transcriptome and miRNA analyses of the response to Corynespora cassiicola in cucumber. Scientific Reports, 2018, 8, 7798.	3.3	43
2	The Two Translationally Controlled Tumor Protein Genes, CsTCTP1 and CsTCTP2, Are Negative Modulators in the Cucumis sativus Defense Response to Sphaerotheca fuliginea. Frontiers in Plant Science, 2018, 9, 544.	3.6	21
3	Comparative proteomic analysis of cucumber roots infected by Fusarium oxysporum f. sp. cucumerium Owen. Physiological and Molecular Plant Pathology, 2016, 96, 77-84.	2.5	17
4	Nucleotide-Binding Leucine-Rich Repeat Genes CsRSF1 and CsRSF2 Are Positive Modulators in the Cucumis sativus Defense Response to Sphaerotheca fuliginea. International Journal of Molecular Sciences, 2021, 22, 3986.	4.1	15
5	Proteome-level investigation of Cucumis sativus-derived resistance to Sphaerotheca fuliginea. Acta Physiologiae Plantarum, 2014, 36, 1781-1791.	2.1	12
6	Cucumber Mildew Resistance Locus O Interacts with Calmodulin and Regulates Plant Cell Death Associated with Plant Immunity. International Journal of Molecular Sciences, 2019, 20, 2995.	4.1	11
7	Genetic diversity of diazotrophs and total bacteria in the phyllosphere of Pyrus serotina, Prunus armeniaca, Prunus avium, and Vitis vinifera. Canadian Journal of Microbiology, 2019, 65, 642-652.	1.7	10
8	Regulation of Growth and Salt Resistance in Cucumber Seedlings by Hydrogen-Rich Water. Journal of Plant Growth Regulation, 2023, 42, 134-153.	5.1	10
9	A comparative cell wall proteomic analysis of cucumber leaves under Sphaerotheca fuliginea stress. Acta Physiologiae Plantarum, 2016, 38, 1.	2.1	9
10	Mildew Resistance Locus O Genes CsMLO1 and CsMLO2 Are Negative Modulators of the Cucumis sativus Defense Response to Corynespora cassiicola. International Journal of Molecular Sciences, 2019, 20, 4793.	4.1	9
11	Functional Identification of Corynespora cassiicola-Responsive miRNAs and Their Targets in Cucumber. Frontiers in Plant Science, 2019, 10, 668.	3.6	9
12	Molecular characterization, expression analysis and heterologous expression of two translationally controlled tumor protein genes from Cucumis sativus. PLoS ONE, 2017, 12, e0184872.	2.5	7
13	Proteomic analysis of Cucumis sativus cotyledons after glucohexaose treatment as a part of ROS accumulation related resistance mechanism. Proteome Science, 2014, 12, 34.	1.7	5
14	Photosynthesis Regulation by Glucohexaose Through Redox Changes in Cucumis sativus. Journal of Plant Growth Regulation, 2014, 33, 571-578.	5.1	2
15	Draft Genome Sequence of Paenibacillus polymyxa EBL06, a Plant Growth-Promoting Bacterium Isolated from Wheat Phyllosphere. Genome Announcements, 2015, 3, .	0.8	1